



PCT

RAW SEQUENCE LISTING

DATE: 07/08/2004

PATENT APPLICATION: US/10/500,346

TIME: 15:12:53

Input Set : A:\3190-059-sequence-listing.txt

Output Set: N:\CRF4\07082004\J500346.raw

3 <110> APPLICANT: ENDO, Yaeta

4 SAWASAKI, Tatsuya

6 <120> TITLE OF INVENTION: Nucleotidic Sequence Having Translation Efficiency

Regulatory Activity

and Use Thereof

E--> 9 <131> 3190-059

W--> 0 <130> FILE REFERENCE: OK

C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/500,346

12 <141> CURRENT FILING DATE: 2004-06-25

14 <150> PRIOR APPLICATION NUMBER: PCT/JP02/13756

15 <151> PRIOR FILING DATE: 2002-12-27

17 <150> PRIOR APPLICATION NUMBER: JP 2001-396941

18 <151> PRIOR FILING DATE: 2001-12-27

20 <160> NUMBER OF SEQ ID NOS: 145

22 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
Corrected Diskette Needed

(pg.1) ↻

ERRORED SEQUENCES

1787 <210> SEQ ID NO: 145

1788 <211> LENGTH: 33

1789 <212> TYPE: DNA

1790 <213> ORGANISM: Artificial Sequence

1792 <220> FEATURE:

1793 <221> NAME/KEY: misc_feature

1794 <223> OTHER INFORMATION: Synthetic DNA.

1796 <400> SEQUENCE: 145

1797 cgatttaggt gacactatag aactcaccat etc

33

E--> 1801 1/44

E--> 1803 1

E--> 1805 11/43

E--> 1806 Page 32

E--> 1808 Page 31

E--> 1810 Page 2

RAW SEQUENCE LISTING ERROR SUMMARY
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27
Seq#:28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51
Seq#:52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75
Seq#:76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99
Seq#:100,101,102,103,104,105,106,107,108,109,110,111,112,113,114,115,116,117
Seq#:118,119,120,121,122,123,124,125,126,127,128,129,130,131,132,133,134,135
Seq#:136,137,138,139,140,141,142,143,144

VERIFICATION SUMMARY

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Input Set : A:\3190-059-sequence-listing.txt

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L:9 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:58 M:283 W: Missing Blank Line separator, <400> field identifier
L:417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:1801 M:254 E: No. of Bases conflict, LENGTH:Input:44 Counted:34 SEQ:145
L:1801 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=145
L:1805 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:1806 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1808 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1810 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1810 M:252 E: No. of Seq. differs, <211> LENGTH:Input:33 Found:47 SEQ:145